EP0807675A2

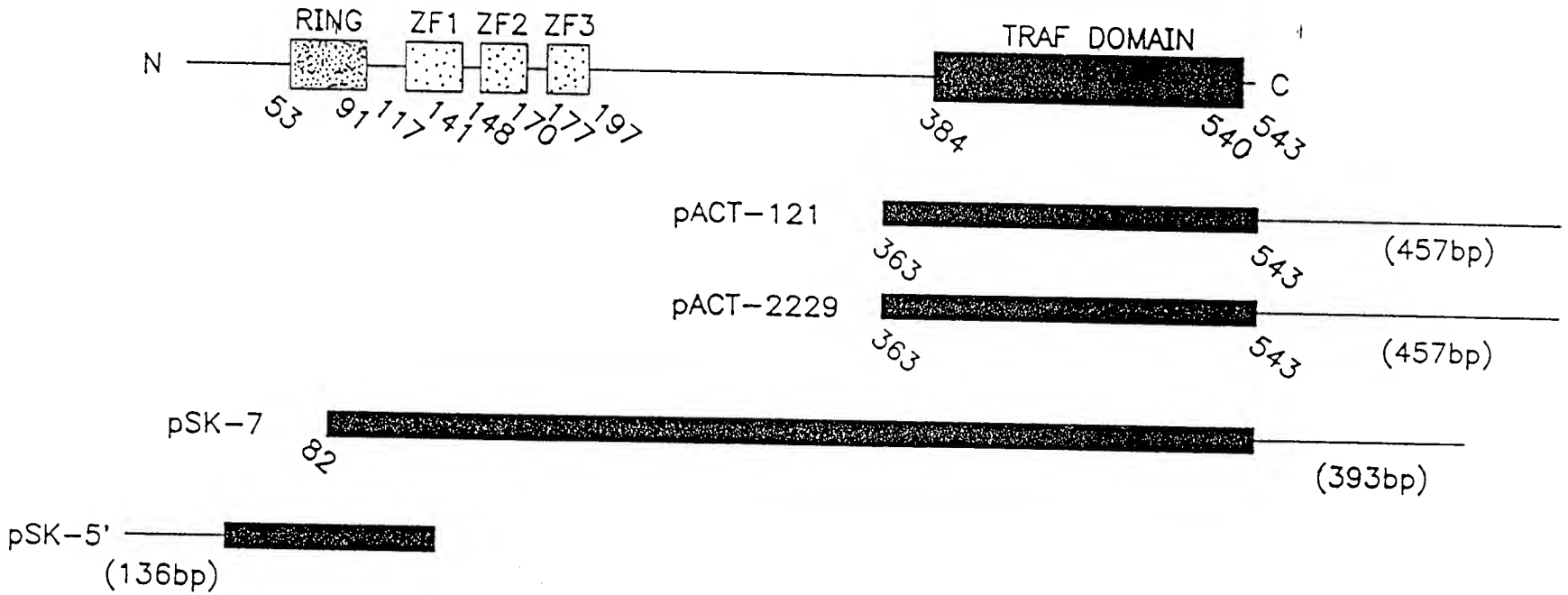


FIG. 2

EP 0 803 570 A2

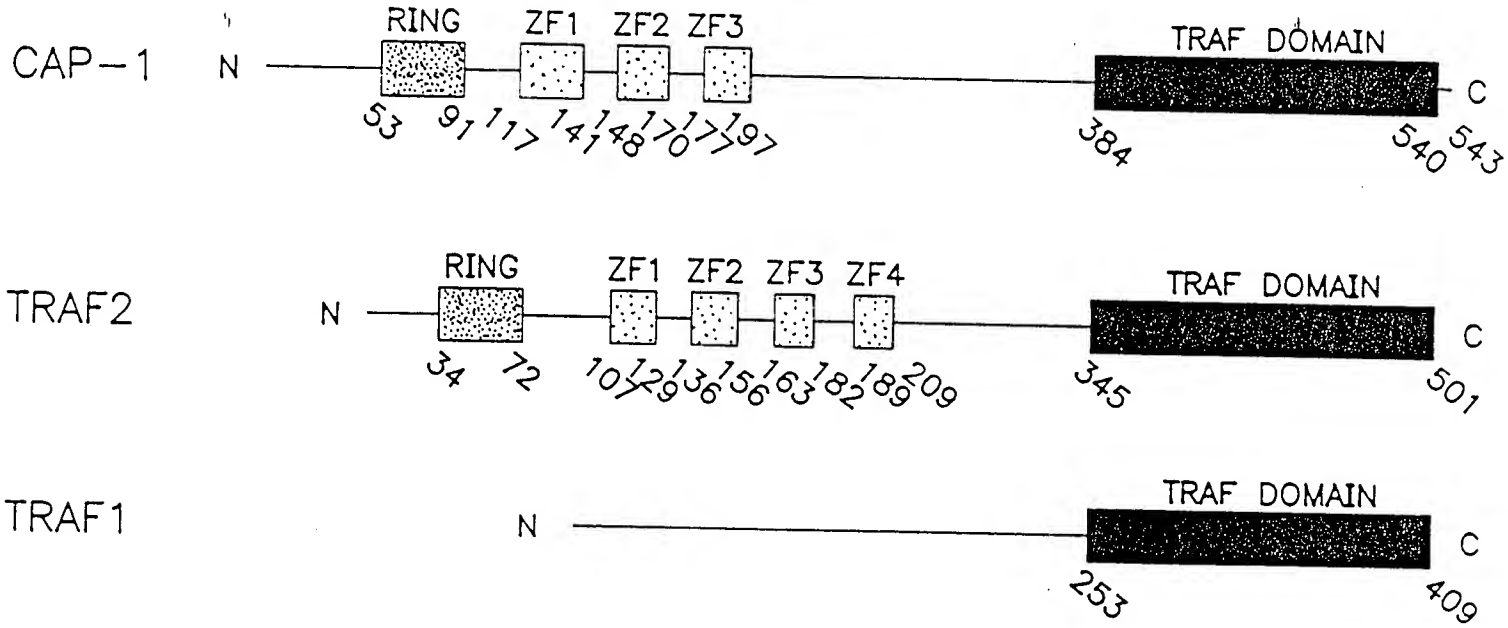
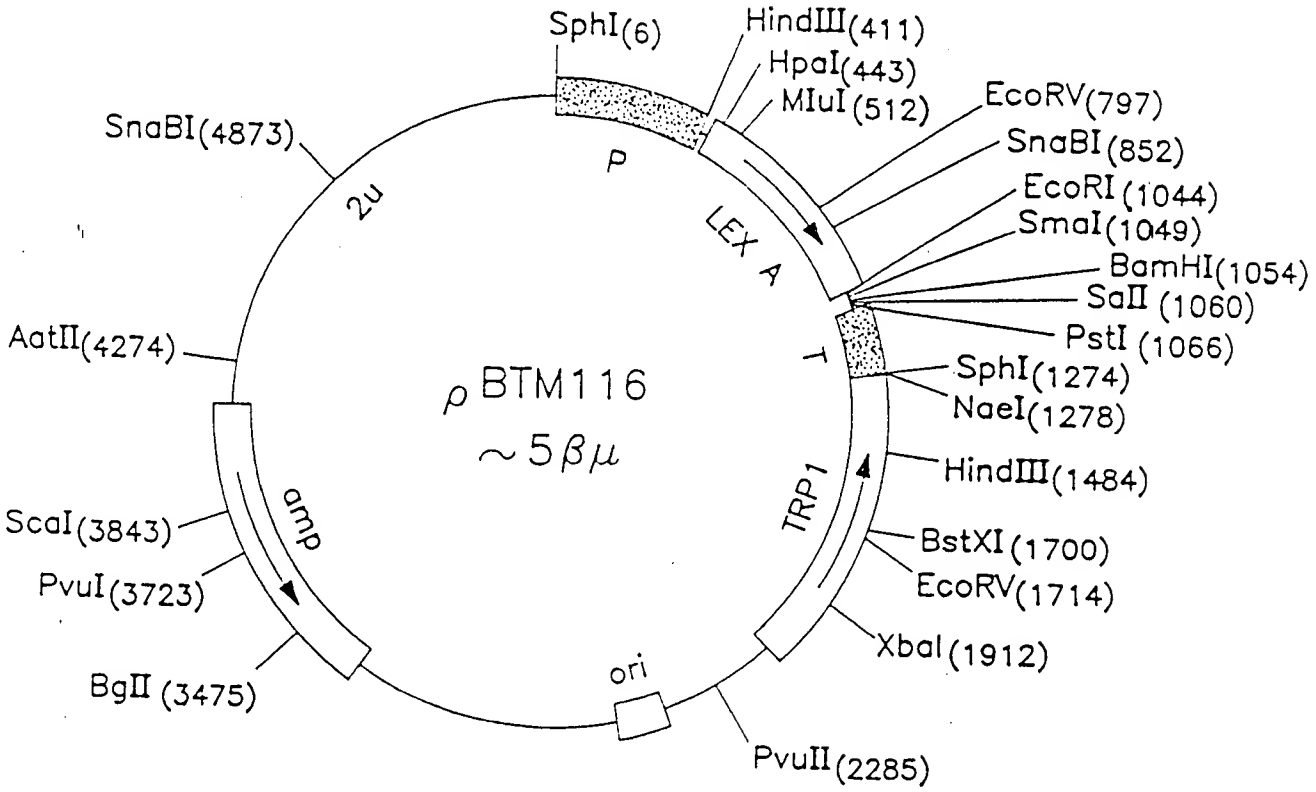


FIG. 3

EDGETO: PHO2560



POLYLINKER READING FRAME:

GAA TTC CCG GGG ATC CGT CGA CCT GCA G

EcoRI SmaI BamHI SalI PstI

FIG. 4

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-136                                                                    AGAAGTGATGCCACTTGGTTAAGGTCCCAGAGCAGGTCAGAATCAGACCTAGGATCA  -80
 -79 GAAACCTGGCTCCTGGCTCCTGGCTCCCTACTCTTCTAAGGATCGCTGTCCTGACAGAAGAGAACTCCTCTTTCCTAAA  -1

   1 ATG GAG TCG AGT AAA AAG ATG GAC TCT CCT GGC GCG CTG CAG ACT AAC CCG CCG CTA AAG   60
   1 M   E   S   S   K   K   M   D   S   P   G   A   L   Q   T   N   P   P   L   K    20

  61 CTG CAC ACT GAC CGC AGT GCT GGG ACG CCA GTT TTT GTC CCT GAA CAA GGA GGT TAC AAG  120
  21 L   H   T   D   R   S   A   G   T   P   V   F   V   P   E   Q   G   G   Y   K    40

 121 GAA AAG TTT GTG AAG ACC GTG GAG GAC AAG TAC AAG TGT GAG AAG TGC CAC CTG GTG CTG  180
  41 E   K   F   V   K   T   V   E   D   K   Y   K   C   E   K   C   H   L   V   L    60

 181 TGC AGC CCG AAG CAG ACC GAG TGT GGG CAC CGC TCC TGC GAG AGC TGC ATG GCG GCC CTG  240
  61 C   S   P   K   Q   T   E   C   G   H   R   S   C   E   S   C   M   A   A   L    80

 241 CTG AGC TCT TCA AGT CCA AAA TGT ACA GCG TGT CAA GAG AGC ATC GTT AAA GAT AAG GTG  300
  81 L   S   S   S   S   P   K   C   T   A   C   Q   E   S   I   V   K   D   K   V   100

 301 TTT AAG GAT AAT TGC TGC AAG AGA GAA ATT CTG GCT CTT CAG ATC TAT TGT CGG AAT GAA  360
 101 F   K   D   N   C   C   K   R   E   I   L   A   L   Q   I   Y   C   R   N   E   120

 361 AGC AGA GGT TGT GCA GAG CAG TTA ACG CTG GGA CAT CTG CTG GTG CAT TTA AAA AAT GAT  420
 121 S   R   G   C   A   E   Q   L   T   L   G   H   L   L   V   H   L   K   N   D   140

 421 TGC CAT TTT GAA GAA CTT CCA TGT GTG CGT CCT GAC TGC AAA GAA AAG GTC TTG AGG AAA  480
 141 C   H   F   E   E   L   P   C   V   R   P   D   C   K   E   K   V   L   R   K   160

 481 GAC CTG CGA GAC CAC GTG GAG AAG GCG TGT AAA TAC CGG GAA GCC ACA TGC AGC CAC TGC  540
 161 D   L   R   D   H   V   E   K   A   C   K   Y   R   E   A   T   C   S   H   C   180

 541 AAG AGT CAG GTT CCG ATG ATC GCG CTG CAG AAA CAC GAA GAC ACC GAC TGT CCC TGC GTG  600
 181 K   S   Q   V   P   M   I   A   L   Q   K   H   E   D   T   D   C   P   C   V   200
```

FIG. 1A

```
601 GTG GTG TCC TGC CCT CAC AAG TGC AGC GTC CAG ACT CTC CTG AGG AGC GAG GGG ACA AAC  660
201 V   V   S   C   P   H   K   C   S   V   Q   T   L   L   R   S   E   G   T   N    220

661 CAG CAG ATC AAG GCC CAC GAG GCC AGC TCC GCC GTG CAG CAC GTC AAC CTG CTG AAG GAG  720
221 Q   Q   I   K   A   H   E   A   S   S   A   V   Q   H   V   N   L   L   K   E    240

721 TGG AGC AAC TCG CTC GAA AAG AAG GTT TCC TTG TTG CAG AAT GAA AGT GTA GAA AAA AAC  780
241 W   S   N   S   L   E   K   K   V   S   L   L   Q   N   E   S   V   E   K   N    260

781 AAG AGC ATA CAA AGT TTG CAC AAT CAG ATA TGT AGC TTT GAA ATT GAA ATT GAG AGA CAA  840
261 K   S   I   Q   S   L   H   N   Q   I   C   S   F   E   I   E   I   E   R   Q    280

841 AAG GAA ATG CTT CGA AAT AAT GAA TCC AAA ATC CTT CAT TTA CAG CGA GTG ATA GAC AGC  900
281 K   E   M   L   R   N   N   E   S   K   I   L   H   L   Q   R   V   I   D   S    300

901 CAA GCA GAG AAA CTG AAG GAG CTT GAC AAG GAG ATC CGG TCC TTC CGG CAG AAC TGG GAG  960
301 Q   A   E   K   L   K   E   L   D   K   E   I   R   S   F   R   Q   N   W   E    320

961 GAA GCA GAC AGC ATG AAG AGC AGC GTG GAG TCC CTC CAG AAC CGC GTG ACC GAG CTG GAG 1020
321 E   A   D   S   M   K   S   S   V   E   S   L   Q   N   R   V   T   E   L   E    340

1021 AGC GTG GAC AAG AGC GCG GGG CAA GTG GCT CGG AAC ACA GGC CTG CTG GAG TCC CAG CTG 1080
 341 S   V   D   K   S   A   G   Q   V   A   R   N   T   G   L   L   E   S   Q   L    360

1081 AGC CGG CAT GAC CAG ATG CTG AGT GTG CAC GAC ATC CGC CTA GCC GAC ATG GAC CTG CGC 1140
 361 S   R   H   D   Q   M   L   S   V   H   D   I   R   L   A   D   M   D   L   R    380

1141 TTC CAG GTC CTG GAG ACC GCC AGC TAC AAT GGA GTG CTC ATC TGG AAG ATT CGC GAC TAC 1200
 381 F   Q   V   L   E   T   A   S   Y   N   G   V   L   I   W   K   I   R   D   Y    400
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FIG. 1B

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1201 AAG CGG CGG AAG CAG GAG GCC GTC ATG GGG AAG ACC CTG TCC CTT TAC AGC CAG CCT TTC  1260
 401 K   R   R   K   Q   E   A   V   M   G   K   T   L   S   L   Y   S   Q   P   F    420

1261 TAC ACT GGT TAC TTT GGC TAT AAG ATG TGT GCC AGG GTC TAC CTG AAC GGG GAC GGG ATG  1320
 421 Y   T   G   Y   F   G   Y   K   M   C   A   R   V   Y   L   N   G   D   G   M    440

1321 GGG AAG GGG ACG CAC TTG TCG CTG TTT TTT GTC ATC ATG CGT GGA GAA TAT GAT GCC CTG  1380
 441 G   K   G   T   H   L   S   L   F   F   V   I   M   R   G   E   Y   D   A   L    460

1381 CTT CCT TGG CCG TTT AAG CAG AAA GTG ACA CTC ATG CTG ATG GAT CAG GGG TCC TCT CGA  1440
 461 L   P   W   P   F   K   Q   K   V   T   L   M   L   M   D   Q   G   S   S   R    480

1441 CGT CAT TTG GGA GAT GCA TTC AAG CCC GAC CCC AAC AGC AGC AGC TTC AAG AAG CCC ACT  1500
 481 R   H   L   G   D   A   F   K   P   D   P   N   S   S   S   F   K   K   P   T    500

1501 GGA GAG ATG AAT ATC GCC TCT GGC TGC CCA GTC TTT GTG GCC CAA ACT GTT CTA GAA AAT  1560
 501 G   E   M   N   I   A   S   G   C   P   V   F   V   A   Q   T   V   L   E   N    520

1561 GGG ACA TAT ATT AAA GAT GAT ACA ATT TTT ATT AAA GTC ATA GTG GAT ACT TCG GAT CTG  1620
 521 G   T   Y   I   K   D   D   T   I   F   I   K   V   I   V   D   T   S   D   L    540

1621 CCC GAT CCC TGATAAGTAGCTGGGGAGGTGGATTTAGCAGAAGGCAACTCCTCTGGGGGATTTGAACCGGTCTGTC  1696
 541 P   D   P                                                                       543

1697 TTCACTGAGGTCCTCGCGCTCAGAAAAGGACCTTGTGAGACGGAGGAAGCGGCAGAAGGCGGACGCGTGCCGGCGGGAG  1775
1776 GAGCCACGCGTGAGACACCTGACACGTTTTATAATAGACTAGCCACACTTCACTCTGAAGAATTATTTATCCTTCAACA  1854
1855 AGCATAAATATTGCTGTCAGAGAAGGTTTTCATTTTCATTTTTAAAGATCTAGTTAATTAAGGTGGAAAACATATATGC  1933
1934 TAAACAAAAGAAACATGATTTTTCTTCCTTAAACTTGAACACCAAAAAACACACACACACACACACACGTGGGGATAGC  2012
2013 TGGACATGTCAGCATGTTAAGTAAAAGGAGAATTTATGAAATAGTAATGCAATTCTGATATCTTCTTTCTAAAATTCAA  2091
2092 GAGTGCAATTTTG                                                                    2104
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FIG. IC